

Kam



1600

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/839,577A

DATE: 05/09/2003
TIME: 08:48:50

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF4\05092003\I839577A.raw

ENTERED

4 <110> APPLICANT: Hahn, Klaus M.
 5 Toutchkine, Alexei
 6 Muthyalu, Rajeev
 7 Kraynov, Vadim
 8 Burton, Dennis R.
 9 Chamberlain, Chester
 10 The Scripps Research Institute et al.
 12 <120> TITLE OF INVENTION: Labeled Peptides, Proteins and Antibodies and Processes and
 Intermediates
 13 Useful for their Preparation
 15 <130> FILE REFERENCE: 1361.007US1
 17 <140> CURRENT APPLICATION NUMBER: US 09/839,577A
 18 <141> CURRENT FILING DATE: 2001-04-20
 20 <150> PRIOR APPLICATION NUMBER: US 60/279,302
 21 <151> PRIOR FILING DATE: 2001-03-28
 23 <150> PRIOR APPLICATION NUMBER: PCT/US00/26821
 24 <151> PRIOR FILING DATE: 2000-09-29
 26 <150> PRIOR APPLICATION NUMBER: US 60/218,113
 27 <151> PRIOR FILING DATE: 2000-07-13
 29 <160> NUMBER OF SEQ ID NOS: 15
 31 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 33 <210> SEQ ID NO: 1
 34 <211> LENGTH: 44
 35 <212> TYPE: PRT
 36 <213> ORGANISM: Homo sapiens
 38 <400> SEQUENCE: 1
 39 Lys Lys Lys Glu Lys Glu Arg Pro Glu Ile Ser Leu Pro Ser Asp Phe
 40 1 5 10 15
 41 Glu His Thr Ile His Val Gly Phe Asp Ala Cys Thr Gly Glu Phe Thr
 42 20 25 30
 43 Gly Met Pro Glu Gln Trp Ala Arg Leu Leu Gln Thr
 44 35 40
 46 <210> SEQ ID NO: 2
 47 <211> LENGTH: 16
 48 <212> TYPE: PRT
 49 <213> ORGANISM: Artificial Sequence
 51 <220> FEATURE:
 52 <223> OTHER INFORMATION: A synthetic peptide.
 54 <400> SEQUENCE: 2
 55 Ala Lys Ala Ala Arg Ala Ala Ala Lys Ala Ala Arg Ala Cys Ala
 56 1 5 10 15
 58 <210> SEQ ID NO: 3
 59 <211> LENGTH: 6
 60 <212> TYPE: PRT

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61 <213> ORGANISM: Artificial Sequence
 63 <220> FEATURE:
 64 <223> OTHER INFORMATION: A synthetic peptide.
 W--> 66 <221> NAME/KEY: SITE
 67 <222> LOCATION: 3
 68 <223> OTHER INFORMATION: Xaa = SAOD = alpha-Boc-beta[N-(2-Chlorobenzylloxycarbonyl)-N-Methylaminoxy Acetyl]-alpha,beta-Diaminopropionic Acid [Boc-2-Cl-Z-(SA)Dapa-OH].
 W--> 72 <221> SITE
 73 <222> LOCATION: 6
 74 <223> OTHER INFORMATION: Xaa = MPAL = The C-terminal mercaptopropionyl-leucine group generated by cleavage of a peptide from TAMPAL resin.
 W--> 77 <400> 3
 W--> 78 Leu Tyr Xaa Ala Gly Xaa
 79 1 5
 81 <210> SEQ ID NO: 4
 82 <211> LENGTH: 5
 83 <212> TYPE: PRT
 84 <213> ORGANISM: Artificial Sequence
 86 <220> FEATURE:
 87 <223> OTHER INFORMATION: A synthetic peptide.
 89 <400> SEQUENCE: 4
 90 Cys Arg Ala Asn Lys
 91 1 5
 93 <210> SEQ ID NO: 5
 94 <211> LENGTH: 10
 95 <212> TYPE: PRT
 96 <213> ORGANISM: Artificial Sequence
 98 <220> FEATURE:
 99 <223> OTHER INFORMATION: A synthetic peptide.
 W--> 101 <221> NAME/KEY: SITE
 102 <222> LOCATION: 3
 103 <223> OTHER INFORMATION: Xaa = SAOD = alpha-Boc-beta[N-(2-Chlorobenzylloxycarbonyl)-N-Methylaminoxy Acetyl]-alpha,beta-Diaminopropionic Acid [Boc-2-Cl-Z-(SA)Dapa-OH]
 W--> 107 <400> 5
 W--> 108 Leu Tyr Xaa Ala Gly Cys Arg Ala Asn Lys
 109 1 5 10
 111 <210> SEQ ID NO: 6
 112 <211> LENGTH: 33
 113 <212> TYPE: PRT
 114 <213> ORGANISM: Artificial Sequence
 116 <220> FEATURE:
 117 <223> OTHER INFORMATION: A synthetic peptide.
 119 <400> SEQUENCE: 6
 120 Cys Glu Tyr Arg Ile Asp Arg Val Arg Leu Phe Val Asp Lys Leu Asp
 121 1 5 10 15
 122 Asn Ile Ala Gln Val Pro Arg Val Gly Ala Ala His His His His His
 123 20 25 30

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124 His
 127 <210> SEQ ID NO: 7
 128 <211> LENGTH: 33
 129 <212> TYPE: PRT
 130 <213> ORGANISM: Artificial Sequence
 132 <220> FEATURE:
 133 <223> OTHER INFORMATION: A synthetic peptide.
 135 <400> SEQUENCE: 7
 136 Cys Glu Tyr Arg Ile Asp Arg Val Leu Phe Val Asp Lys Leu Asp
 137 1 5 10 15
 138 Asn Ile Ala Gln Val Pro Arg Val Gly Ala Ala His His His His
 139 20 25 30
 140 His
 143 <210> SEQ ID NO: 8
 144 <211> LENGTH: 28
 145 <212> TYPE: PRT
 146 <213> ORGANISM: Artificial Sequence
 148 <220> FEATURE:
 149 <223> OTHER INFORMATION: A synthetic peptide.
 W--> 151 <221> NAME/KEY: SITE
 152 <222> LOCATION: 1
 153 <223> OTHER INFORMATION: Xaa = SAOD = alpha-Boc-beta[N-(2-Chlorobenzylloxycarbonyl)-N-
 154 Methylaminoxy
 155 Acetyl]-alpha,beta-Diaminopropionic Acid [Boc-2-Cl-Z-(SA)Dapa-OH]
 W--> 157 <221> SITE
 158 <222> LOCATION: 28
 159 <223> OTHER INFORMATION: Xaa = MPAL = The C-terminal mercaptopropionyl-leucine group
 generated by
 160 cleavage of a peptide from TAMPAL resin.
 W--> 162 <400> 8
 W--> 163 Xaa Lys Lys Lys Glu Lys Glu Arg Pro Glu Ile Ser Leu Pro Ser Asp
 164 1 5 10 15
 165 Phe Glu His Thr Ile His Val Gly Phe Asp Ala Xaa
 166 20 25
 168 <210> SEQ ID NO: 9
 169 <211> LENGTH: 18
 170 <212> TYPE: PRT
 171 <213> ORGANISM: Homo sapiens
 173 <400> SEQUENCE: 9
 174 Cys Thr Gly Glu Phe Thr Gly Met Pro Glu Gln Trp Ala Arg Leu Leu
 175 1 5 10 15
 176 Gln Thr
 179 <210> SEQ ID NO: 10
 180 <211> LENGTH: 10
 181 <212> TYPE: PRT
 182 <213> ORGANISM: Artificial Sequence
 184 <220> FEATURE:
 185 <223> OTHER INFORMATION: A synthetic peptide.
 W--> 187 <221> NAME/KEY: SITE
 188 <222> LOCATION: 3

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189 <223> OTHER INFORMATION: Xaa = SAOD = alpha-Boc-beta[N-(2-Chlorobenzylloxycarbonyl)-N-Methylaminoxy Acetyl]-alpha,beta-Diaminopropionic Acid [Boc-2-Cl-Z-(SA)Dapa-OH]

W--> 192 <400> 10

W--> 193 Leu Tyr Xaa Ala Gly Cys Arg Ala Asn Lys

194	1	5	10
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196 <210> SEQ ID NO: 11

197 <211> LENGTH: 45

198 <212> TYPE: PRT

199 <213> ORGANISM: Artificial Sequence

201 <220> FEATURE:

202 <223> OTHER INFORMATION: A synthetic peptide.

W--> 204 <221> NAME/KEY: SITE

205 <222> LOCATION: 1

206 <223> OTHER INFORMATION: Xaa = SAOD = alpha-Boc-beta[N-(2-Chlorobenzylloxycarbonyl)-N-Methylaminoxy Acetyl]-alpha,beta-Diaminopropionic Acid [Boc-2-Cl-Z-(SA)Dapa-OH]

W--> 209 <400> 11

W--> 210 Xaa Lys Lys Lys Glu Lys Glu Arg Pro Glu Ile Ser Leu Pro Ser Asp

211	1	5	10	15
-----	---	---	----	----

212 Phe Glu His Thr Ile His Val Gly Phe Asp Ala Cys Thr Gly Glu Phe

213	20	25	30
-----	----	----	----

214 Thr Gly Met Pro Glu Gln Trp Ala Arg Leu Leu Gln Thr

215	35	40	45
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217 <210> SEQ ID NO: 12

218 <211> LENGTH: 38

219 <212> TYPE: PRT

220 <213> ORGANISM: Artificial Sequence

222 <220> FEATURE:

223 <223> OTHER INFORMATION: A synthetic peptide.

W--> 225 <221> NAME/KEY: SITE

226 <222> LOCATION: 3

227 <223> OTHER INFORMATION: Xaa = SAOD = alpha-Boc-beta[N-(2-Chlorobenzylloxycarbonyl)-N-Methylaminoxy Acetyl]-alpha,beta-Diaminopropionic Acid [Boc-2-Cl-Z-(SA)Dapa-OH]

W--> 230 <400> 12

W--> 231 Leu Tyr Xaa Ala Gly Cys Glu Tyr Arg Ile Asp Arg Val Arg Leu Phe

232	1	5	10	15
-----	---	---	----	----

233 Val Asp Lys Leu Asp Asn Ile Ala Gln Val Pro Arg Val Gly Ala Ala

234	20	25	30
-----	----	----	----

235 His His His His His

236	35
-----	----

238 <210> SEQ ID NO: 13

239 <211> LENGTH: 120

240 <212> TYPE: PRT

241 <213> ORGANISM: Homo sapiens

243 <400> SEQUENCE: 13

244 Asp Ile Gln Asn Pro Asp Ile Thr Ser Ser Arg Tyr Arg Gly Leu Pro

245	1	5	10	15
-----	---	---	----	----

246 Ala Pro Gly Pro Ser Pro Ala Asp Lys Lys Arg Ser Gly Lys Lys Lys

247	20	25	30
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248 Ile Ser Lys Ala Asp Ile Gly Ala Pro Ser Gly Phe Lys His Val Ser

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249 35 40 45
 250 His Val Gly Trp Asp Pro Gln Asn Gly Phe Asp Val Asn Asn Leu Asp
 251 50 55 60
 252 Pro Asp Leu Arg Ser Leu Phe Ser Arg Ala Gly Ile Ser Glu Ala Gln
 253 65 70 75 80
 254 Leu Thr Asp Ala Glu Thr Ser Lys Leu Ile Tyr Asp Phe Ile Glu Asp
 255 85 90 95
 256 Gln Gly Gly Leu Glu Ala Val Arg Gln Glu Met Arg Arg Gln Glu Pro
 257 100 105 110
 258 Leu Pro Pro Pro Pro Pro Ser
 259 115 120
 262 <210> SEQ ID NO: 14
 263 <211> LENGTH: 502
 264 <212> TYPE: PRT
 265 <213> ORGANISM: Homo sapiens
 267 <400> SEQUENCE: 14
 268 Met Ser Gly Gly Pro Met Gly Gly Arg Pro Gly Gly Arg Gly Ala Pro
 269 1 5 10 15
 270 Ala Val Gln Gln Asn Ile Pro Ser Thr Leu Leu Gln Asp His Glu Asn
 271 20 25 30
 272 Gln Arg Leu Phe Glu Met Leu Gly Arg Lys Cys Leu Thr Leu Ala Thr
 273 35 40 45
 274 Ala Val Val Gln Leu Tyr Leu Ala Leu Pro Pro Gly Ala Glu His Trp
 275 50 55 60
 276 Thr Lys Glu His Cys Gly Ala Val Cys Phe Val Lys Asp Asn Pro Gln
 277 65 70 75 80
 278 Lys Ser Tyr Phe Ile Arg Leu Tyr Gly Leu Gln Ala Gly Arg Leu Leu
 279 85 90 95
 280 Trp Glu Gln Glu Leu Tyr Ser Gln Leu Val Tyr Ser Thr Pro Thr Pro
 281 100 105 110
 282 Phe Phe His Thr Phe Ala Gly Asp Asp Cys Gln Ala Gly Leu Asn Phe
 283 115 120 125
 284 Ala Asp Glu Asp Glu Ala Gln Ala Phe Arg Ala Leu Val Gln Glu Lys
 285 130 135 140
 286 Ile Gln Lys Arg Asn Gln Arg Gln Ser Gly Asp Arg Arg Gln Leu Pro
 287 145 150 155 160
 288 Pro Pro Pro Thr Pro Ala Asn Glu Glu Arg Arg Gly Gly Leu Pro Pro
 289 165 170 175
 290 Leu Pro Leu His Pro Gly Gly Asp Gln Gly Gly Pro Pro Val Gly Pro
 291 180 185 190
 292 Leu Ser Leu Gly Leu Ala Thr Val Asp Ile Gln Asn Pro Asp Ile Thr
 293 195 200 205
 294 Ser Ser Arg Tyr Arg Gly Leu Pro Ala Pro Gly Pro Ser Pro Ala Asp
 295 210 215 220
 296 Lys Lys Arg Ser Gly Lys Lys Ile Ser Lys Ala Asp Ile Gly Ala
 297 225 230 235 240
 298 Pro Ser Gly Phe Lys His Val Ser His Val Gly Trp Asp Pro Gln Asn
 299 245 250 255
 300 Gly Phe Asp Val Asn Asn Leu Asp Pro Asp Leu Arg Ser Leu Phe Ser

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; Xaa Pos. 3,6
Seq#:5; Xaa Pos. 3
Seq#:8; Xaa Pos. 1,28
Seq#:10; Xaa Pos. 3
Seq#:11; Xaa Pos. 1
Seq#:12; Xaa Pos. 3

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq# :10; Line(s) 189
Seq# :11; Line(s) 206
Seq# :12; Line(s) 227

VERIFICATION SUMMARY

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Input Set : A:\PTO.AMC.txt

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L:66 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:72 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:77 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:3
L:78 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
L:101 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:107 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:5
L:108 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
L:151 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:157 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
L:162 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:8
L:163 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
M:341 Repeated in SeqNo=8
L:187 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:192 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:10
L:193 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:204 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:209 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:11
L:210 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
L:225 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:230 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:12
L:231 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0